In the Claims

Cancel claims 4-22 and 28-41, without prejudice.

RECEIVED

Please amend claims 1-3 and 23 to read as follows:

JUN 0 8 2001

1. AMENDED An oligonucleotide comprising at least one concatenation coding for a polypertide 1600/2900 with formula $(P-K)_n$, where:

n is a whole number of 2 or more;

P represents a proline amino acid residue;

K represents a lysine amino acid residue;

the symbol "-" represents a bond between the two amino acid residues, in particular a peptide type bond, the n (P-K) units also being bonded together by such bonds, for example peptide type bonds.

- 2. The oligonucleotide according to claim 1, comprising a concatenation coding for a polypeptide with formula (P-K)_n where n is a whole number of 3 or more, and preferably n is equal to 4, 5, 6, 7, 8, 9, 10 or 151.
- The oligonucleotide according to claim 1, comprising a concatenation coding for a polypeptide with formula (P-K)_n, in which the sequence of n (P-K) units is interrupted by one or more amino acid residues other than P or K residues.

23. A cloning and/or expression vector, which is one of plasmids pP20yZ (CNCM N° I-1640), pH30yZ or pH45yZ (CNCM N° I-1639).

Add claims 42-82 as follows:

42. An oligonucleotide comprising at least one concatenation coding for a

polypeptide with formula (P-K)_n, where:

n is a whole number of 2 or more;

P represents a proline amino acid residue;

K represents a lysine amino acid residue;

the symbol "-"represents a bond between the two amino acid residues, in particular a peptide type bond, the n (P-K) units also being bonded together by such bonds, for example peptide type bonds.

- 43. The oligonucleotide according to claim 42, comprising a concatenation coding for a polypeptide with formula (P-K)_n where n is a whole number of 3 or more, and preferably n is equal to 4, 5, 6, 7, 8, 9, 10 or 15.
- 44. The oligonucleotide according to claim 42, comprising a concatenation coding for a polypeptide with formula (P-K)_n, in which the sequence of n (P-K) units is interrupted by one or more amino acid residues other than P or K residues.

45. The oligonucleotide according to claim 42, wherein the concatenation coding for the polypeptide comprising the n (P-K) units is completed at its 5' end and/or at its 3' end by one or more codons coding, for example, for at least one lysine residue at the N-terminal extremity of the formed polypeptide.

- 46. The oligonucleotide according to claim 45, which codes for a polypeptide with formula (P-K), formula K-(P-K)₄, or with formula 2K(P-K)₄.
- 47. A recombinant nucleotide sequence comprising a concatenation of nucleotides coding for a plant protein reserve, which further comprises an oligonucleotide according to any one of claims 42 to 46, inserted at one site of the nucleotide concatenation selected such that:

expression of the nucleotide sequence in a particular plant cell enables a modified protein reserve to be produced which is localized in that cell in a manner identical to or similar to the normal protein reserve which would be expressed in the same cell under the same conditions by the corresponding normal coding nucleotide concatenation; and/or

153 Const the modified protein reserve coded by the recombinant nucleotide sequence is immunologically recognized by antibodies produced against the corresponding normal protein reserve.

- 48. The nucleotide sequence according to claim 47, wherein the coding nucleotide concatenation codes for a protein reserve which is naturally low in lysine.
- 49. The nucleotide sequence according to claim 48, wherein the coding nucleotide concatenation codes for a protein reserve naturally produced by a plant for use in animal or human foodstuffs.
- 50. The nucleotide sequence according to claim 48, wherein the coding nucleotide concatenation codes for a protein reserve naturally produced by a plant from the cereal family.
- 51. The nucleotide sequence according to claim 48, wherein the coding nucleotide concatenation codes for a protein reserve naturally produced by a plant from the legume or crucifer family.
- 52. The nucleotide sequence according to claim 50, wherein the coding nucleotide concatenation codes for a maize protein reserve.
- 53. The nucleotide sequence according to claim 52, wherein the coding nucleotide concatenation codes for a protein reserve from the zein family.
- 54. The nucleotide sequence according to claim 53, wherein the coding nucleotide concatenation codes for a protein reserve which is maize γ -zein.
- 55. The nucleotide sequence according to claim 54, wherein the nucleotide concatenation coding for the maize γ-zein has the sequence shown in Figure 9.

- The nucleotide sequence according to claim 48, wherein the coding nucleotide concatenation codes for a protein reserve of a plant selected from the following: soya, sunflower, tobacco, wheat, oats, alfalfa, rice, oilseed rape, sorghum, and *Arabidopsis thaliana*.
- 57. The nucleotide sequence according to claim 47, wherein the protein reserve encoded by the coding nucleotide concatenation is maize γ-zein, and wherein the oligonucleotide is inserted in place of or following a Pro-X domain or in a Pro-X domain naturally present in the maize γ-zein.
- 58. A recombinant nucleotide sequence, which comprises a nucleotide sequence according to claim 47 under the control of an expression promoter.
- 59. The recombinant nucleotide sequence according to claim 58, wherein the promoter is a specific promoter for a given cell tissue, for example a promoter which is specific for expression in grains, and/or in the leaves of plants.
- 60. The nucleotide sequence according to claim 58, wherein the expression promoter is that of maize γ -zein.
- 61. The nucleotide sequence according to claim 58, wherein the expression promoter is the promoter CaMV35S.
- 62. The nucleotide sequence according to claim 57, which codes for one of the polypeptides P20γZ or H45γZ with the sequences shown in Figures 11 and 10, respectively.
- 63. A cloning and/or expression vector, which comprises, at a site which is not essential for replication, a nucleotide sequence in accordance with claim 47.

- 64. A cloning and/or expression vector, which is one of plasmids pP20yZ (CNCM N° I-1640), pH30yZ or pH45yZ (CNCM N° I-1639).
- 65. A polypeptide coded by a sequence according to claim 47.
- 66. A lysine-enriched modified maize γ -zein, which is coded by a nucleotide sequence according to claim 54.
- 67. A lysine-enriched modified maize γ -zein, the amino acid sequence of which is modified by at least one polypeptide with formula $(P-K)_n$ or with formula $2K(P-K)_n$, where:

n is a whole number of 2 or more;

P represents a proline amino acid residue;

K represents a lysine amino acid residue;

the symbol "-" represents a bond between the two amino acid residues, in particular a peptide type bond, the n (P-K) units being bonded together by bonds, in particular peptide type bonds, said polypeptide having formula $(P-K)_n$ or $K-(P-K)_n$ being substituted for a sequence naturally present in the normal maize γ -zein or being inserted with deletion of one or more amino acids of the amino acid sequence for normal maize γ -zein, or being added to the normal γ -zein amino acid sequence, the insertion site for the polypeptide being selected such that:

when the modified lysine-rich γ -zein is produced in a host cell, in particular in a plant cell, it is localized in identical or similar manner to the normal maize γ -zein which would be produced under the same conditions in the same host cell; and/or

the modified maize γ -zein is recognized by antibodies directed against the normal maize γ -zein.

B3

- The modified maize y-zein according to claim 67, which is the protein P20yZ or 68. the protein H30yZ or the protein H45yZ.
- A recombinant host cell, which comprises a nucleotide sequence according to 69. claim 47.
- The host cell according to claim 69, which is a bacterium, for example E. coli or 70. Agrobacterium tumefaciens.
- 71. The host cell according to claim 69, which is a plant cell.
- The host cell according to claim 71, which is a plant seed cell. 72.
- 73. The host cell according to claim 72, which is a cell from maize seed endosperm.
- 74. The host cell according to claim 73, which contains a nucleotide sequence according to claim 54, integrated in its genome in a stable manner.
- The host cell according to claim 73, which produces a lysine-enriched modified 75. maize y-zein according to claim 67.
- 76. The host cell according to claim 71, which is a soya, sunflower, tobacco, wheat, oats, alfalfa, rice, oilseed rape, sorghum or Arabidopsis cell.
- 77. Seeds producing a polypeptide according to any one of claims 65 to 68.
- A plant producing a polypeptide according to any one of claims 65 to 68. 78.
- 79. The plant according to claim 78, which is a maize plant.
- Seeds obtained from plants according to claim 78.

mat (S)

- 81. A method of producing plants or seeds expressing a modified protein reserve, which comprises the steps of:
 - a) transforming a plant cell with a nucleotide sequence according to claim 47, or a vector according to claim 63, under conditions enabling the modified protein reserve coded by the nucleotide sequence to be expressed in a stable and functional manner;
 - b) regenerating plants from the plant cell transformed in step a), to obtain plants expressing the modified protein reserve;
 - c) if necessary, obtaining seeds from the modified plants obtained in step b).
- 82. The method according to claim 81, wherein the plant is maize and the protein reserve is γ -zein.